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[Sequence Listing]
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<120> Humanized anti-CD47 antibody
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cagtgaaggt ttc 133
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gcagtctcag atc 133
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 gtc cac tcc cag gtg cag ctg gtg cag tct ggg gct gag gtg aag aag 96
 Val His Ser Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys
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Ala Asn His Val Ile His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu
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gag tgg atg gga tat att tat cct tac aat gat ggt act aag tat aat 240
Glu Trp Met Gly Tyr Ile Tyr Pro Tyr Asn Asp Gly Thr Lys Tyr Asn
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gag aag ttc aag gac aga gtc acg atg acc cgg gac acg tcc acg agc 288
Glu Lys Phe Lys Asp Arg Val Thr Met Thr Arg Asp Thr Ser Thr Ser
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aca gtc tac atg gag ttg agc agt ctc aga tct gag gac acg gcc gtc 336
Thr Val Tyr Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val
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                              85
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tat tat tgt gct aga ggg ggt tac tat act tac gac gac tgg ggc caa 384
Tyr Tyr Cys Ala Arg Gly Gly Tyr Tyr Thr Tyr Asp Asp Trp Gly Gln
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gtc cac tcc cag gtg cag ctg gtg cag tct ggg gct gag gtg aag aag 96
Val His Ser Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys
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cct ggg gcc tca gtg aag gtt tcc tgt aag gca tct gga tac acc ttc 144
Pro Gly Ala Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe
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gcc aac cat gtt att cac tgg gtg cga cag gcc cct gga caa ggg ctt 192
Ala Asn His Val Ile His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu
                                          40
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gag tgg atg gga tat att tat cct tac aat gat ggt act aag tat aat 240
Glu Trp Met Gly Tyr Ile Tyr Pro Tyr Asn Asp Gly Thr Lys Tyr Asn
                                                          60
                                      55
                  50
gag aag tic aag gac aga gic acg aig acc ica gac acg icc acg agc 288
Glu Lys Phe Lys Asp Arg Val Thr Met Thr Ser Asp Thr Ser Thr Ser
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                                                      75
              65
aca gic tac atg gag tig agc agt cic aga tot gag gac acg gcc gtc 336
Thr Val Tyr Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val
                                                 . 90
          80
tat tat tgt gct aga ggg ggt tac tat act tac gac gac tgg ggc caa 384
Tyr Tyr Cys Ala Arg Gly Gly Tyr Tyr Thr Tyr Asp Asp Trp Gly Gln
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gga acc ctg gtc acc gtc tcc tca ggt gag tgg atc cgc g 424
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gtc cac tcc cag gtg cag ctg gtg cag tct ggg gct gag gtg aag aag 96
Val His Ser Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys
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cct ggg gcc tca gtg aag gtt tcc tgt aag gca tct gga tac acc ttc 144
Pro Gly Ala Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe
                         20
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acc aac cat git att cac igg gig cga cag gcc cci gga caa ggg cii 192
Thr Asn His Val Ile His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu
                                         40
 30
                     35
                                                             45
gag tgg atg gga tat att tat cct tac aat gat ggt act aag tat aat 240
Glu Trp Met Gly Tyr Ile Tyr Pro Tyr Asn Asp Gly Thr Lys Tyr Asn
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gag aag ttc aag gac aga gtc acg atg acc tca gac acg tcc acg agc 288
Glu Lys Phe Lys Asp Arg Val Thr Met Thr Ser Asp Thr Ser Thr Ser
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aca gtc tac atg gag ttg agc agt ctc aga tct gag gac acg gcc gtc 336
Thr Val Tyr Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val
                            85
         80
tat tat tgt gct aga ggg ggt tac tat act tac gac 'gac tgg ggc caa 384
Tyr Tyr Cys Ala Arg Gly Gly Tyr Tyr Thr Tyr Asp Asp Trp Gly Gln
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                        100
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gga acc ctg gtc acc gtc tcc tca ggt gag tgg atc cgc g 424
Gly Thr Leu Val Thr Val Ser Ser
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<213 Mouse, Human
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gtc cac tcc cag gtg cag ctg gtg cag tct ggg gct gag gtg aag aag 96
Val His Ser Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys
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cct ggg gcc tca gtg aag gtt tcc tgt aag gca tct gga tac acc ttc 144
Pro Gly Ala Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe
                                             25
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gcc aac cat gtt att cac tgg gtg cga cag gcc cct gga caa ggg ctt 192
Ala Asn His Val Ile His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu
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gag igg aig gga tat att tai cci tac aai gai ggi aci aag tai aai 240
Glu Trp Met Gly Tyr Ile Tyr Pro Tyr Asn Asp Gly Thr Lys Tyr Asn
                                      55
                 50
gag aag ttc aag gac aaa gtc acg atg acc tca gac acg tcc acg agc 288
Glu Lys Phe Lys Asp Lys Val Thr Met Thr Ser Asp Thr Ser Thr Ser
                                                      75
                                  70
             65
aca gtc tac atg gag ttg agc agt ctc aga tct gag gac acg gcc gtc 336
Thr Val Tyr Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val
                                                  90
                              85
         80
tat tat tgt gct aga ggg ggt tac tat act tac gac gac tgg ggc caa 384
Tyr Tyr Cys Ala Arg Gly Gly Tyr Tyr Thr Tyr Asp Asp Trp Gly Gln
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gga acc ctg gtc acc gtc tcc tca ggt gag tgg atc cgc g 424
Gly Thr Leu Val Thr Val Ser Ser
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<212> DNA
<213> Mouse, Human
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gic cac tcc cag gig cag cig gig cag ici ggg gci gag gig aag aag
Val His Ser Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys
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cct ggg gcc tca gtg aag gtt tcc tgt aag gca tct gga tac acc ttc 144
Pro Gly Ala Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe
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gcc aac cat gtt att cac tgg gtg cga cag gcc cct gga caa ggg ctt 192
Ala Asn His Val Ile His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu
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gag tgg atg gga tat att tat cct tac aat gat ggt act aag tat aat 240
Glu Trp Met Gly Tyr Ile Tyr Pro Tyr Asn Asp Gly Thr Lys Tyr Asn
                                       55
                  50
gag aag ttc aag gac aga gtc acg ctg acc tca gac acg tcc acg agc 288
Glu Lys Phe Lys Asp Arg Val Thr Leu Thr Ser Asp Thr Ser Thr Ser
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 aca gtc tac atg gag tig agc agt ctc aga tct gag gac acg gcc gtc 336
 Thr Val Tyr Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val
                               85
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 tat tat tgt gct aga ggg ggt tac tat act tac gac gac tgg ggc caa 384
 Tyr Tyr Cys Ala Arg Gly Gly Tyr Tyr Thr Tyr Asp Asp Trp Gly Gln
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 gga acc ctg gtc acc gtc tcc tca ggt gag tgg atc cgc g 424
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<400> 21
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<213 Mouse, Human
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gic cac tcc cag gig cag cig gig cag ict ggg gci gag gig aag aag 96
Val His Ser Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys
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 cct ggg gcc tca gtg aag gtt tcc tgt aag gca tct gga tac acc tic 144
Pro Gly Ala Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe
                                              25
                          20
 acc aac cat git att cac igg gig cga cag gcc cci gga caa ggg cii 192
 Thr Asn His Val Ile His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu
                                                               45
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gag tgg atg gga tat att tat cct tac aat gat ggt act aag tat aat 240
Glu Trp Met Gly Tyr Ile Tyr Pro Tyr Asn Asp Gly Thr Lys Tyr Asn
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                                     55
gag aag ttc aag gac aga gtc acg atg acc tca gac acg tcc acg agc 288
Glu Lys Phe Lys Asp Arg Val Thr Met Thr Ser Asp Thr Ser Thr Ser
                                                     75
                                 70
             65
aca gic tac atg gag tig agc agt cic aga tot gac gac acg gcc gic 336
Thr Val Tyr Met Glu Leu Ser Ser Leu Arg Ser Asp Asp Thr Ala Val
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tat tat igi gci aga ggg ggi tac tat act tac gac gac igg ggc caa 384
Tyr Tyr Cys Ala Arg Gly Gly Tyr Tyr Thr Tyr Asp Asp Trp Gly Gln
                                             105
                        100
     95
gga acc ctg gtc acc gtc tcc tca ggt gag tgg atc cgc g 424
Gly Thr Leu Val Thr Val Ser Ser
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                     115
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ccagtgaata acatggtt 18
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<211> 49
<212> DNA
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<213> Mouse, Human
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gic cac tcc cag gig cag cig gig cag tci ggg gct gag gig aag aag
Val His Ser Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys
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              1
cct ggg gcc tca gtg cag gtt tcc tgt aag gca tct gga tac acc ttc 144
Pro Gly Ala Ser Val Gln Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe
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     15
acc aac cat git att cac tgg cig cga cag gcc cct gga caa ggg cii 192
Thr Asn His Val Ile His Trp Leu Arg Gln Ala Pro Gly Gln Gly Leu
                      35
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gag tgg atg gga tat att tat cct tac aat gat ggt act aag tat aat 240
Glu Trp Met Gly Tyr Ile Tyr Pro Tyr Asn Asp Gly Thr Lys Tyr Asn
                                                           60
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                  50
gag aag tic aag gac aga gtc acg atg acc tca gac acg tcc atc agc 288
Glu Lys Phe Lys Asp Arg Val Thr Met Thr Ser Asp Thr Ser Ile Ser
                                                       75
                                   70
              65
 aca gcc tac atg gag ttg agc agt ctc aga tct gac gac acg gcc gtc 336
 Thr Ala Tyr Met Glu Leu Ser Ser Leu Arg Ser Asp Asp Thr Ala Val
                              85
          80
 tat tat tgt gct aga ggg ggt tac tat act tac gac gac tgg ggc caa 384
 Tyr Tyr Cys Ala Arg Gly Gly Tyr Tyr Thr Tyr Asp Asp Trp Gly Gln
                                              105
                          100
      95
 gca acc ctg gtc acc gtc tcc tca ggt gag tgg atc cgc g 424
 Ala Thr Leu Val Thr Val Ser Ser
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 110
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 ggc tcc agt ggg gat git gtg atg act cag tct cca ctc tcc ctg ccc 96
 Gly Ser Ser Gly Asp Val Val Met Thr Gln Ser Pro Leu Ser Leu Pro
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 gtc acc ctt gga cag ccg gcc tcc atc tcc tgc aga tca agt cag agc 144
 Val Thr Leu Gly Gln Pro Ala Ser Ile Ser Cys Arg Ser Ser Gln Ser
                                                   25
                               20
          15
 ctt gig cac agt aat gga aag acc tat tta cat igg tit cag cag agg 192 .
 Leu Val His Ser Asn Gly Lys Thr Tyr Leu His Trp Phe Gln Gln Arg
                           35
      30
 cca ggc caa ici cca agg cgc cta att tat aaa git icc aac cga iii 240
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Pro Gly Gln Ser Pro Arg Arg Leu Ile Tyr Lys Val Ser Asn Arg Phe
                                          55
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 45
tct ggt gtc cca gac aga ttc agc ggc agt ggg tca ggc act gat ttc 288
Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe
                                                          75
                                      70
                  65
aca ctg aaa atc agc agg gtg gag gct gag gat gtt gga gtt tat tac 336
Thr Leu Lys Ile Ser Arg Val Glu Ala Glu Asp Val Gly Val Tyr Tyr
tgc tct caa agt aca cat gtt ccg tac acg ttt ggc cag ggg acc aag 384
Cys Ser Gln Ser Thr His Val Pro Tyr Thr Phe Gly Gln Gly Thr Lys
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                             100
          95
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Leu Glu Ile Lys
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 <211> 412
 <212> DNA
 <213> Mouse, Human
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<400> 40

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ggc tcc agt ggg gat gtt gtg atg act cag tct cca ctc tcc ctg ccc 96
Gly Ser Ser Gly Asp Val Val Met Thr Gln Ser Pro Leu Ser Leu Pro
                                                      10
                                   5
             -1
                  1
gtc acc ctt gga cag ccg gcc tcc atc tcc tgc aga tca agt cag agc 144
Val Thr Leu Gly Gln Pro Ala Ser Ile Ser Cys Arg Ser Ser Gln Ser
                              20
         15
cti gig cac agi aai gga aag acc tai ita cai igg tii cag cag agg 192
Leu Val His Ser Asn Gly Lys Thr Tyr Leu His Trp Phe Gln Gln Arg
                                              40
                          35
     `30
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Pro Gly Gln Ser Pro Arg Leu Leu lle Tyr Lys Val Ser Asn Arg Phe
                                                               60
                                          55
                      50
 45
tct ggt gtc cca gac aga ttc agc ggc agt ggg tca ggc act gat ttc 288
Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe
                                      70
                  65
aca cig aaa atc agc agg gtg gag gct gag gat gtt gga gtt tat tac 336
 Thr Leu Lys Ile Ser Arg Val Glu Ala Glu Asp Val Gly Val Tyr Tyr
                                                       90
                                  85
              80
 tgc tct caa agt aca cat gtt ccg tac acg ttt ggc cag ggg acc aag 384
 Cys Ser Gln Ser Thr His Val Pro Tyr Thr Phe Gly Gln Gly Thr Lys
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          95
 ctg gag atc aaa cgt gag tgg atc cgc g 412
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<400> 43
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Met Arg Leu Pro Ala Gln Leu Leu Gly Leu Leu Met Leu Trp Val Pro
ggc tcc agt ggg gat gtt gtg atg act cag tct cca ctc tcc ctg ccc 96
Gly Ser Ser Gly Asp Val Val Met Thr Gln Ser Pro Leu Ser Leu Pro
                                                      10
                                   5
             -1
gtc acc ctt gga cag ccg gcc tcc atc tcc tgc aga tca agt cag agc 144
Val Thr Leu Gly Gln Pro Ala Ser Ile Ser Cys Arg Ser Ser Gln Ser
                              20
          15
ctt gtg cac agt aat gga aag acc tat tta cat tgg ttt cag cag agg 192
Leu Val His Ser Asn Gly Lys Thr Tyr Leu His Trp Phe Gln Gln Arg
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      30
 cca ggc caa tct cca agg cgc cta att tat aaa gtt tcc aac cga ttt 240
 Pro Gly Gln Ser Pro Arg Arg Leu Ile Tyr Lys Val Ser Asn Arg Phe
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 Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe
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 aca cig aaa atc agc agg gig gag gct gag gat git gga git tat iic 336
 Thr Leu Lys Ile Ser Arg Val Glu Ala Glu Asp Val Gly Val Tyr Phe
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                                   85
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 tgc tct caa agt aca cat gtt ccg tac acg ttt ggc cag ggg acc aag 384
 Cys Ser Gln Ser Thr His Val Pro Tyr Thr Phe Gly Gln Gly Thr Lys
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 ggc tcc agt ggg gat gtt gtg atg act cag tct cca ctc tcc ctg ccc 96
· Gly Ser Ser Gly Asp Val Val Met Thr Gln Ser Pro Leu Ser Leu Pro
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 gtc acc ctt gga cag ccg gcc tcc atc tcc tgc aga tca agt cag agc 144
 Val Thr Leu Gly Gln Pro Ala Ser Ile Ser Cys Arg Ser Ser Gln Ser
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 ctt gtg cac agt aat gga aag acc tat tta cat tgg tac cag cag agg 192
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Leu Val His Ser Asn Gly Lys Thr Tyr Leu His Trp Tyr Gln Gln Arg
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Pro Gly Gln Ser Pro Arg Arg Leu Ile Tyr Lys Val Ser Asn Arg Phe
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tct ggt gtc cca gac aga tic agc ggc agt ggg ica ggc act gat tic 288
Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe
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aca ctg aaa atc agc agg gtg gag gct gag gat gtt gga gtt tat tac 336
Thr Leu Lys Ile Ser Arg Val Glu Ala Glu Asp Val Gly Val Tyr Tyr
                                                      90
                                  85
             80
tgc tct caa agt aca cat gtt ccg tac acg ttt ggc cag ggg acc aag 384
Cys Ser Gln Ser Thr His Val Pro Tyr Thr Phe Gly Gln Gly Thr Lys
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ctg gag atc aaa cgt gag tgg atc cgc g 412
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<211> 412

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ggc tcc agt ggg gat git gtg atg act cag tct cca ctc tcc ctg ccc 96
Gly Ser Ser Gly Asp Val Val Met Thr Gln Ser Pro Leu Ser Leu Pro
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gtc acc ctt gga cag ccg gcc tcc atc tcc tgc aga tca agt cag agc 144
Val Thr Leu Gly Gln Pro Ala Ser Ile Ser Cys Arg Ser Ser Gln Ser
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ctt gig cac agi aai gga aag acc tat tia cai igg tii cig cag agg 192
Leu Val His Ser Asn Gly Lys Thr Tyr Leu His Trp Phe Leu Gln Arg
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Pro Gly Gln Ser Pro Arg Arg Leu Ile Tyr Lys Val Ser Asn Arg Phe
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Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe
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 Thr Leu Lys Ile Ser Arg Val Glu Ala Glu Asp Val Gly Val Tyr Tyr
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                                  85
              80
 tgc tct caa agt aca cat gtt ccg tac acg ttt ggc cag ggg acc aag 384
 Cys Ser Gln Ser Thr His Val Pro Tyr Thr Phe Gly Gln Gly Thr Lys
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ggc tcc agt ggg gat gtt gtg atg act cag tct cca ctc tcc ctg ccc 96
Gly Ser Ser Gly Asp Val Val Met Thr Gln Ser Pro Leu Ser Leu Pro
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Val Thr Leu Gly Gln Pro Ala Ser Ile Ser Cys Arg Ser Ser Gln Ser
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ctt gtg cac agt aat gga aag acc tat tta cat tgg tat ctg cag aag 192
Leu Val His Ser Asn Gly Lys Thr Tyr Leu His Trp Tyr Leu Gln Lys
                          35
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cca ggc cag tct cca aga ctc ctg atc tac aaa gtt tcc aac cga ttt 240
Pro Gly Gln Ser Pro Arg Leu Leu Ile Tyr Lys Val Ser Asn Arg Phe
                                                               60
                                          55
                      50
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 tct ggt gtc cca gac aga ttc agc ggc agt ggg tca ggc act gat ttc 288
 Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe
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                  65
 aca ctg aaa atc agc agg gtg gag gct gag gat gtt gga gtt tat tac 336
 Thr Leu Lys Ile Ser Arg Val Glu Ala Glu Asp Val Gly Val Tyr Tyr
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90
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tgc tct caa agt aca cat gtt ccg tac acg ttt ggc cag ggg acc aag 384
Cys Ser Gln Ser Thr His Val Pro Tyr Thr Phe Gly Gln Gly Thr Lys
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ctg gag atc aaa cgt gag tgg atc cgc g 412
Leu Glu Ile Lys
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ggc tcc agt ggg gat att gtg atg act cag tct cca ctc tcc ctg ccc 96
Gly Ser Ser Gly Asp Ile Val Met Thr Gln Ser Pro Leu Ser Leu Pro
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gic acc cct gga gag ccg gcc tcc atc tcc tgc aga tca agt cag agc 144
Val Thr Pro Gly Glu Pro Ala Ser Ile Ser Cys Arg Ser Ser Gln Ser
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ctt gig cac agt aat gga aag acc tat tia cat igg tat cig cag aag 192
Leu Val His Ser Asn Gly Lys Thr Tyr Leu His Trp Tyr Leu Gln Lys
                                              40
                          35
      30
cca ggc cag tct cca aga ctc ctg atc tac aaa gtt tcc aac cga ttt 240
Pro Gly Gln Ser Pro Arg Leu Leu Ile Tyr Lys Val Ser Asn Arg Phe
                                                              60
                                          55
                      50
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 tct ggt gtc cca gac aga ttc agc ggc agt ggg tca ggc act gat ttc 288
Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe
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                  65
 aca cig aaa atc agc agg gig gag gct gat gat git gga att tat tac 336
 Thr Leu Lys Ile Ser Arg Val Glu Ala Asp Asp Val Gly Ile Tyr Tyr
                                                       90
                                   85
              80
 tgc tct caa agt aca cat gtt ccg tac acg ttt ggc cag ggg acc aag 384
 Cys Ser Gln Ser Thr His Val Pro Tyr Thr Phe Gly Gln Gly Thr Lys
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                              100
 ctg gag atc aaa cgt gag tgg atc cgc g 412
 Leu Glu Ile Lys
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 Val His Ser Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys
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 Pro Gly Ala Ser Val Gln Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe
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  Thr Asn His Val Met His Trp Leu Arg Gln Ala Pro Gly Gln Gly Leu
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  gag igg aig gga tat ait tat cct tac aat gat ggt act aag tat aat 240
  Glu Trp Met Gly Tyr Ile Tyr Pro Tyr Asn Asp Gly Thr Lys Tyr Asn
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                    50
  gag aag tic aag ggc aga gtc acg atg acc tca gac acg tcc atc agc 288
  Glu Lys Phe Lys Gly Arg Val Thr Met Thr Ser Asp Thr Ser Ile Ser
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tat tat tgt gct aga ggg ggt tac tat tct tac gac gac tgg ggc caa 384
Tyr Tyr Cys Ala Arg Gly Gly Tyr Tyr Ser Tyr Asp Asp Trp Gly Gln
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 ggc tcc agt ggg gat att gtg atg act cag tct cca ctc tcc ctg ccc 96
 Gly Ser Ser Gly Asp Ile Val Met Thr Gln Ser Pro Leu Ser Leu Pro
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Val Thr Pro Gly Glu Pro Ala Ser Ile Ser Cys Arg Ser Ser Gln Ser
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cit cig cac agi aag gga aac acc tat tia cag igg tat cig cag aag 192
Leu Leu His Ser Lys Gly Asn Thr Tyr Leu Gln Trp Tyr Leu Gln Lys
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cca ggc cag tct cca aga ctc ctg atc tac aaa gtt tcc aac cga ttt 240
Pro Gly Gln Ser Pro Arg Leu Leu Ile Tyr Lys Val Ser Asn Arg Phe
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tct ggt gtc cca gac aga ttc agc ggc agt ggg tca ggc act gat ttc 288
Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe
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                  65
aca ctg aaa atc agc agg gtg gag gct gat gat gtt gga att tat tac 336
Thr Leu Lys Ile Ser Arg Val Glu Ala Asp Asp Val Gly Ile Tyr Tyr
                                                      90
                                  85
              80
tgc tct caa agt aca cat gtt ccg tac acg ttt ggc cag ggg acc aag 384
Cys Ser Gln Ser Thr His Val Pro Tyr Thr Phe Gly Gln Gly Thr Lys
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Gly Gly Gly Ser
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gic gac tcc cag gig cag cig gig cag ici ggg gci gag gig aag aag 96
Val Asp Ser Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys
cct ggg gcc tca gtg cag gtt tcc tgt aag gca tct gga tac acc ttc 144
Pro Gly Ala Ser Val Gln Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe
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                         20
     15
acc aac cat git ait cac tgg ctg cga cag gcc cct gga caa ggg cit 192
Thr Asn His Val Ile His Trp Leu Arg Gln Ala Pro Gly Gln Gly Leu
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gag tgg atg gga tat att tat cct tac aat gat ggt act aag tat aat 240
Glu Trp Met Gly Tyr Ile Tyr Pro Tyr Asn Asp Gly Thr Lys Tyr Asn
                                      55
                  50
gag aag tic aag gac aga gic acg atg acc tca gac acg icc atc agc 288
Glu Lys Phe Lys Asp Arg Val Thr Met Thr Ser Asp Thr Ser Ile Ser
                                                      75
                                  70
              65
aca gcc tac atg gag ttg agc agt ctc aga tct gac gac acg gcc gtc 336
Thr Ala Tyr Met Glu Leu Ser Ser Leu Arg Ser Asp Asp Thr Ala Val
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          80
 tat tat tgt gct aga ggg ggt tac tat act tac gac gac tgg ggc caa 384
Tyr Tyr Cys Ala Arg Gly Gly Tyr Tyr Thr Tyr Asp Asp Trp Gly Gln
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 Ala Thr Leu Val Thr Val Ser Ser Gly Gly Gly Ser Asp Ile Val
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                                          120
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 Ser Ile Ser Cys Arg Ser Ser Gln Ser Leu Val His Ser Asn Gly Lys
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 Thr Tyr Leu His Trp Tyr Leu Gln Lys Pro Gly Gln Ser Pro Arg Leu
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 ctg atc tac aaa git tcc aac cga tit tct ggt gic cca gac aga tic 624
 Leu Ile Tyr Lys Val Ser Asn Arg Phe Ser Gly Val Pro Asp Arg Phe
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85

80

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Tyr Tyr Cys Ala Arg Gly Gly Tyr Tyr Ser Tyr Asp Asp Trp Gly Gln
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gca acc ctg gtc aca gtc tcg agt ggt ggc gga ggt tcc gat att gtg 432
Ala Thr Leu Val Thr Val Ser Ser Gly Gly Gly Gly Ser Asp Ile Val
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atg act cag tot cca ctc tcc ctg ccc gtc acc cct gga gag ccg gcc 480
Met Thr Gln Ser Pro Leu Ser Leu Pro Val Thr Pro Gly Glu Pro Ala
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tcc atc tcc tgc aga tca agt cag agc ctt ctg cac agt aag gga aac 528
Ser Ile Ser Cys Arg Ser Ser Gln Ser Leu Leu His Ser Lys Gly Asn
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Thr Tyr Leu Gln Trp Tyr Leu Gln Lys Pro Gly Gln Ser Pro Arg Leu
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cig atc tac aaa git tcc aac cga tii tct ggt gic cca gac aga tic 624
Leu Ile Tyr Lys Val Ser Asn Arg Phe Ser Gly Val Pro Asp Arg Phe
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age gge agt ggg tea gge act gat the aca etg aaa ate age agg gtg 672
Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Lys Ile Ser Arg Val
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Glu Ala Asp Asp Val Gly Ile Tyr Tyr Cys Ser Gln Ser Thr His Val
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ccg tac acg tit ggc cag ggg acc aag ctg gag atc aaa taa tga gcg 768
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<212> DNA
<213 Mouse, Human
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gtc gac tcc cag gtg cag ctg gtg cag tct ggg gct gag gtg aag aag 96
Val Asp Ser Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys
cct ggg gcc tca gtg cag gtt tcc tgt aag gca tct gga tac acc ttc 144
Pro Gly Ala Ser Val Gln Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe
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acc aac cat git att cac igg cig cga cag gcc cct gga caa ggg cit 192
Thr Asn His Val Ile His Trp Leu Arg Gln Ala Pro Gly Gln Gly Leu
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gag tgg atg gga tat att tat cct tac aat gat ggt act aag tat aat 240
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Thr	Ala	Tyr	Met	Glu	Leu	Ser	Ser	Leu	Arg	Ser	Asp		Thr	Ala	Val	
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Tyr	Tyr	Cys	Ala	Arg	Gly		Tyr	Tyr	Thr	Tyr		Asp	Trp	Gly	Gin	
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меі	ınr	GIN	ser	130	Leu	261	Leu	110	135	1111	110	OI,	014	Pro 140	,,, u	
100	atc	100	tar		tca	aøt	cag	agc		gig	cac	agt	aat	gga	aag	528
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561	110	501	145	6				150					155			
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ctg	atc	t ac	aaa	gtt	tcc	aac	cga	ttt	tct	ggt	gtc	cca	gac	aga	ttc	624
Leu	Ile	Tyr	Lys	Val	Ser	Asn	Arg	Phe	Ser	Gly	Val	Pro	Asp	Arg	Phe	
	175					180					185					
														agg		672
Ser	Gly	Ser	Gly	Ser	Gly	Thr	Asp	Phe	Thr			Ile	Ser	Arg		
190					195					200					205	=00
														cat		720
Glu	Ala	Asp	Asp			Ile	Tyr	Tyr			Gln	Ser	Thr	His	vai	
				210				•	215					220	~~+	760
														ggt		
Pro	Туг	Thr			Glr	Gly	Thr			GIU	1 11e	Lys		Gly	Gly	
			225				لسيس	230		. ~~-			235		c f m	816
														cag Cln		
Gly	Ser	. Gly	/ Gly	/ GIS	/ G13	ser	61)	/ GI)	ч	61)	361	GIII	rai	Gln	Leu	

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250
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Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr Asn His Val Ile His Trp
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Leu Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met Gly Tyr Ile Tyr
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Pro Tyr Asn Asp Gly Thr Lys Tyr Asn Glu Lys Phe Lys Asp Arg Val
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Thr Met Thr Ser Asp Thr Ser Ile Ser Thr Ala Tyr Met Glu Leu Ser
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Tyr Tyr Thr Tyr Asp Asp Trp Gly Gln Ala Thr Leu Val Thr Val Ser
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Leu Pro Val Thr Pro Gly Glu Pro Ala Ser Ile Ser Cys Arg Ser Ser
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Gln Ser Leu Val His Ser Asn Gly Lys Thr Tyr Leu His Trp Tyr Leu
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Gln Lys Pro Gly Gln Ser Pro Arg Leu Leu Ile Tyr Lys Val Ser Asn
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                         420
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cga ttt tct ggt gtc cca gac aga ttc agc ggc agt ggg tca ggc act 1392
Arg Phe Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Thr
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445 440 435 430 gat tic aca ctg aaa atc agc agg gtg gag gct gat gat gtt gga att 1440 Asp Phe Thr Leu Lys Ile Ser Arg Val Glu Ala Asp Asp Val Gly Ile 460 450 tat tac tgc tct caa agt aca cat gtt ccg tac acg ttt ggc cag ggg 1488 Tyr Tyr Cys Ser Gln Ser Thr His Val Pro Tyr Thr Phe Gly Gln Gly 475 470 465 acc aag ctg gag atc aaa taa tga gcg 1515 Thr Lys Leu Glu Ile Lys 480 <210> 79 <211> 1515 <212> DNA <213 Mouse, Human <400> 79 atg gga tgg agc tgt atc atc ctc ttc ttg gta gca aca gct aca ggt 48 Met Gly Trp Ser Cys Ile Ile Leu Phe Leu Val Ala Thr Ala Thr Gly gtc gac tcc cag gtg cag ctg gtg cag tct ggg gct gag gtg aag aag 96 Val Asp Ser Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys -11 cct ggg gcc tca gtg cag gtt tcc tgt aag gca tct gga tac acc ttc 144 Pro Gly Ala Ser Val Gln Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe 25 20 15 acc aac cat git atg cac tgg ctg cga cag gcc cct gga caa ggg ctt 192 Thr Asn His Val Met His Trp Leu Arg Gln Ala Pro Gly Gln Gly Leu 45 35 gag tgg atg gga tat att tat cct tac aat gat ggt act aag tat aat 240 Glu Trp Met Gly Tyr Ile Tyr Pro Tyr Asn Asp Gly Thr Lys Tyr Asn 60 55 50 gag aag tic aag ggc aga gtc acg atg acc tca gac acg tcc atc agc 288 Glu Lys Phe Lys Gly Arg Val Thr Met Thr Ser Asp Thr Ser Ile Ser 75 70 65 aca gcc tac atg gag ttg agc agt ctc aga tct gac gac acg gcc gtc 336 Thr Ala Tyr Met Glu Leu Ser Ser Leu Arg Ser Asp Asp Thr Ala Val

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Ala Thr Leu Val Thr Val Ser Ser Gly Gly Gly Ser Asp Ile Val
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Thr Tyr Leu Gln Trp Tyr Leu Gln Lys Pro Gly Gln Ser Pro Arg Leu
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Glu Ala Asp Asp Val Gly Ile Tyr Tyr Cys Ser Gln Ser Thr His Val
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Pro Tyr Thr Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys Gly Gly Gly
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Gly Ser Gly Gly Gly Ser Gly Gly Gly Ser Gln Val Gln Leu
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35

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Gly Tyr Ile Tyr Pro Tyr Asn Asp Gly Thr Lys Tyr Asn Glu Lys Phe
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Lys Asp Arg Val Thr Met Thr Ser Asp Thr Ser Ile Ser Thr Ala Tyr
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Met Glu Leu Ser Ser Leu Arg Ser Asp Asp Thr Ala Val Tyr Tyr Cys
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Ala Arg Gly Gly Tyr Tyr Thr Tyr Asp Asp Trp Gly Gln Ala Thr Leu
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Val Thr Val Ser Ser Gly Gly Gly Gly Ser Asp Ile Val Met Thr Gln
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Ser Pro Leu Ser Leu Pro Val Thr Pro Gly Glu Pro Ala Ser Ile Ser
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Cys Arg Ser Ser Gln Ser Leu Val His Ser Asn Gly Lys Thr Tyr Leu
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                                        155
His Trp Tyr Leu Gln Lys Pro Gly Gln Ser Pro Arg Leu Leu Ile Tyr
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Lys Val Ser Asn Arg Phe Ser Gly Val Pro Asp Arg Phe Ser Gly Ser
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540

600

660

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170

190

Lys Val Ser Asn Arg Phe Ser Gly Val Pro Asp Arg Phe Ser Gly Ser

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Gly Ser Gly Thr Asp Phe Thr Leu Lys Ile Ser Arg Val Glu Ala Asp 195 200 205

Asp Val Gly Ile Tyr Tyr Cys Ser Gln Ser Thr His Val Pro Tyr Thr 210 215 220

Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys